

## **STIC Biotechnology Systems Branch**

### **RAW SEQUENCE LISTING** **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/575,261  
Source: IFWP  
Date Processed by STIC: 04/24/2006

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

**<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>**

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):  
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06



IFWP

## RAW SEQUENCE LISTING

DATE: 04/24/2006

PATENT APPLICATION: US/10/575,261

TIME: 16:15:41

Input Set : A:\seq list.txt

Output Set: N:\CRF4\04242006\J575261.raw

3 &lt;110&gt; APPLICANT: KYOWA HAKKO KOGYO CO., LTD.

5 &lt;120&gt; TITLE OF INVENTION: Fusion protein composition

7 &lt;130&gt; FILE REFERENCE: 11613WO1

C--&gt; 9 &lt;140&gt; CURRENT APPLICATION NUMBER: US/10/575,261

C--&gt; 9 &lt;141&gt; CURRENT FILING DATE: 2006-04-10

9 &lt;150&gt; PRIOR APPLICATION NUMBER: P2003-350158

10 &lt;151&gt; PRIOR FILING DATE: 2003-10-08

12 &lt;160&gt; NUMBER OF SEQ ID NOS: 113

14 &lt;170&gt; SOFTWARE: PatentIn Ver. 2.1

Does Not Comply  
Corrected Diskette Needed  
(PS-1, 2, 4, 5, 6)

## ERRORED SEQUENCES

131 <210> SEQ ID NO: 2  
 132 <211> LENGTH: 376 → found 372  
 133 <212> TYPE: PRT  
 134 <213> ORGANISM: Cricetulus griseus  
 136 <400> SEQUENCE: 2  
 137 Met Ala His Ala Pro Ala Ser Cys Pro Ser Ser Arg Asn Ser Gly Asp  
 138 1 5 10 15  
 140 Gly Asp Lys Gly Lys Pro Arg Lys Val Ala Leu Ile Thr Gly Ile Thr  
 141 20 25 30  
 143 Gly Gln Asp Gly Ser Tyr Leu Ala Glu Phe Leu Leu Glu Lys Gly Tyr  
 144 35 40 45  
 146 Glu Val His Gly Ile Val Arg Arg Ser Ser Ser Phe Asn Thr Gly Arg  
 147 50 55 60  
 149 Ile Glu His Leu Tyr Lys Asn Pro Gln Ala His Ile Glu Gly Asn Met  
 150 65 70 75 80  
 152 Lys Leu His Tyr Gly Asp Leu Thr Asp Ser Thr Cys Leu Val Lys Ile  
 E--> 153 85 85 90 90 95 100  
 155 Ile Asn Glu Val Lys Pro Thr Glu Ile Tyr Asn Leu Gly Ala Gln Ser  
 E--> 156 100 105 105 110 115  
 158 His Val Lys Ile Ser Phe Asp Leu Ala Glu Tyr Thr Ala Asp Val Asp  
 E--> 159 115 120 120 125 125 130  
 161 Gly Val Gly Thr Leu Arg Leu Leu Asp Ala Ile Lys Thr Cys Gly Leu  
 E--> 162 130 135 135 140 140 145  
 164 Ile Asn Ser Val Lys Phe Tyr Gln Ala Ser Thr Ser Glu Leu Tyr Gly  
 E--> 165 145 150 150 155 155 160  
 167 Lys Val Gln Glu Ile Pro Gln Lys Glu Thr Thr Pro Phe Tyr Pro Arg  
 E--> 168 165 165 170 175 175 180  
 170 Ser Pro Tyr Gly Ala Ala Lys Leu Tyr Ala Tyr Trp Ile Val Val Asn  
 E--> 171 180 185 185 190 195  
 173 Phe Arg Glu Ala Tyr Asn Leu Phe Ala Val Asn Gly Ile Leu Phe Asn  
 195 200 205

Invalid  
Amino Acid  
numbering

## RAW SEQUENCE LISTING

DATE: 04/24/2006

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Input Set : A:\seq list.txt

Output Set: N:\CRF4\04242006\J575261.raw

E--> 174                    200                    205                    210  
 176 His Glu Ser Pro Arg Arg Gly Ala Asn Phe Val Thr Arg Lys Ile Ser  
 E--> 177        210 215                    215 220                    220 225  
 179 Arg Ser Val Ala Lys Ile Tyr Leu Gly Gln Leu Glu Cys Phe Ser Leu  
 E--> 180 225 230                    230 235                    235 240                    240  
 182 Gly Asn Leu Asp Ala Lys Arg Asp Trp Gly His Ala Lys Asp Tyr Val  
 E--> 183 245                    245 250                    250 255                    255 260  
 185 Glu Ala Met Trp Leu Met Leu Gln Asn Asp Glu Pro Glu Asp Phe Val  
 E--> 186                    260 265                    265 270                    270 275  
 188 Ile Ala Thr Gly Glu Val His Ser Val Arg Glu Phe Val Glu Lys Ser  
 E--> 189        275 280                    280 285                    285 290  
 191 Phe Met His Ile Gly Lys Thr Ile Val Trp Glu Gly Lys Asn Glu Asn  
 E--> 192        290 295                    295 300                    300 305  
 194 Glu Val Gly Arg Cys Lys Glu Thr Gly Lys Ile His Val Thr Val Asp  
 E--> 195 305 310                    310 315                    315 320                    320  
 197 Leu Lys Tyr Tyr Arg Pro Thr Glu Val Asp Phe Leu Gln Gly Asp Cys  
 E--> 198 325                    325 330                    330 335                    335 340  
 200 Ser Lys Ala Gln Gln Lys Leu Asn Trp Lys Pro Arg Val Ala Phe Asp  
 E--> 201                    340 345                    345 350                    350 355  
 204 Glu Leu Val Arg Glu Met Val Gln Ala Asp Val Glu Leu Met Arg Thr  
 E--> 205        355 360                    360 365                    365 370  
 207 Asn Pro Asn Ala  
 E--> 208        370 375  
 471 <210> SEQ ID NO: 7  
 472 <211> LENGTH: 575  
 473 <212> TYPE: PRT  
 474 <213> ORGANISM: Cricetulus griseus  
 476 <400> SEQUENCE: 7  
 477 Met Arg Ala Trp Thr Gly Ser Trp Arg Trp Ile Met Leu Ile Leu Phe  
 478        1                    5                    10                    15  
 480 Ala Trp Gly Thr Leu Leu Phe Tyr Ile Gly Gly His Leu Val Arg Asp  
 481                    20                    25                    30  
 483 Asn Asp His Pro Asp His Ser Ser Arg Glu Leu Ser Lys Ile Leu Ala  
 484                    35                    40                    45  
 486 Lys Leu Glu Arg Leu Lys Gln Gln Asn Glu Asp Leu Arg Arg Met Ala  
 487                    50                    55                    60  
 489 Glu Ser Leu Arg Ile Pro Glu Gly Pro Ile Asp Gln Gly Thr Ala Thr  
 490                    65                    70                    75                    80  
 492 Gly Arg Val Arg Val Leu Glu Glu Gln Leu Val Lys Ala Lys Glu Gln  
 493                    85                    90                    95  
 495 Ile Glu Asn Tyr Lys Lys Gln Ala Arg Asn Asp Leu Gly Lys Asp His  
 496                    100                    105                    110  
 498 Glu Ile Leu Arg Arg Arg Ile Glu Asn Gly Ala Lys Glu Leu Trp Phe  
 499                    115                    120                    125  
 501 Phe Leu Gln Ser Glu Leu Lys Lys Leu Lys Lys Leu Glu Gly Asn Glu  
 502                    130                    135                    140  
 505 Leu Gln Arg His Ala Asp Glu Ile Leu Leu Asp Leu Gly His His Glu  
 506                    145                    150                    155                    160  
 508 Arg Ser Ile Met Thr Asp Leu Tyr Tyr Leu Ser Gln Thr Asp Gly Ala

*Same Gene*

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/575,261

DATE: 04/24/2006

TIME: 16:15:41

Input Set : A:\seq list.txt

Output Set: N:\CRF4\04242006\J575261.raw

```

509          165          170          175
511 Gly Glu Trp Arg Glu Lys Glu Ala Lys Asp Leu Thr Glu Leu Val Gln
512          180          185          190
514 Arg Arg Ile Thr Tyr Leu Gln Asn Pro Lys Asp Cys Ser Lys Ala Arg
515          195          200          205
517 Lys Leu Val Cys Asn Ile Asn Lys Gly Cys Gly Tyr Gly Cys Gln Leu
518          210          215          220
520 His His Val Val Tyr Cys Phe Met Ile Ala Tyr Gly Thr Gln Arg Thr
521 225          230          235          240
523 Leu Ile Leu Glu Ser Gln Asn Trp Arg Tyr Ala Thr Gly Gly Trp Glu
524          245          250          255
526 Thr Val Phe Arg Pro Val Ser Glu Thr Cys Thr Asp Arg Ser Gly Leu
527          260          265          270
529 Ser Thr Gly His Trp Ser Gly Glu Val Lys Asp Lys Asn Val Gln Val
530          275          280          285
532 Val Glu Leu Pro Ile Val Asp Ser Leu His Pro Arg Pro Pro Tyr Leu
533          290          295          300
535 Pro Leu Ala Val Pro Glu Asp Leu Ala Asp Arg Leu Leu Arg Val His
536 305          310          315          320
538 Gly Asp Pro Ala Val Trp Trp Val Ser Gln Phe Val Lys Tyr Leu Ile
539          325          330          335
541 Arg Pro Gln Pro Trp Leu Glu Arg Glu Ile Glu Glu Thr Thr Lys Lys
542          340          345          350
544 Leu Gly Phe Lys His Pro Val Ile Gly Val His Val Arg Arg Thr Asp
545          355          360          365
547 Lys Val Gly Thr Glu Ala Ala Phe His Pro Ile Glu Glu Tyr Met Val
548          370          375          380
550 His Val Glu Glu His Phe Gln Leu Leu Glu Arg Arg Met Lys Val Asp
551 385          390          395          400
553 Lys Lys Arg Val Tyr Leu Ala Thr Asp Asp Pro Ser Leu Leu Lys Glu
554          405          410          415
556 Ala Lys Thr Lys Tyr Ser Asn Tyr Glu Phe Ile Ser Asp Asn Ser Ile
557          420          425          430
559 Ser Trp Ser Ala Gly Leu His Asn Arg Tyr Thr Glu Asn Ser Leu Arg
560          435          440          445
562 Gly Val Ile Leu Asp Ile His Phe Leu Ser Gln Ala Asp Phe Leu Val
563          450          455          460
565 Cys Thr Phe Ser Ser Gln Val Cys Arg Val Ala Tyr Glu Ile Met Gln
566 465          470          475          480
568 Thr Leu His Pro Asp Ala Ser Ala Asn Phe His Ser Leu Asp Asp Ile
569          485          490          495
571 Tyr Tyr Phe Gly Gly Gln Asn Ala His Asn Gln Ile Ala Val Tyr Pro
572          500          505          510
574 His Gln Pro Arg Thr Lys Glu Glu Ile Pro Met Glu Pro Gly Asp Ile
575          515          520          525
577 Ile Gly Val Ala Gly Asn His Trp Asn Gly Tyr Ser Lys Gly Val Asn
578          530          535          540
580 Arg Lys Leu Gly Lys Thr Gly Leu Tyr Pro Ser Tyr Lys Val Arg Glu
581 545          550          555          560

```

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/575,261

DATE: 04/24/2006

TIME: 16:15:41

Input Set : A:\seq list.txt

Output Set: N:\CRF4\04242006\J575261.raw

583 Lys Ile Glu Thr Val Lys Tyr Pro Thr Tyr Pro Glu Ala Glu Lys  
 E--> 584 565 570 575  
 587 <210> SEQ ID NO: 8  
 588 <211> LENGTH: 575  
 589 <212> TYPE: PRT  
 590 <213> ORGANISM: Mus musculus  
 592 <400> SEQUENCE: 8  
 593 Met Arg Ala Trp Thr Gly Ser Trp Arg Trp Ile Met Leu Ile Leu Phe  
 594 1 5 10 15  
 596 Ala Trp Gly Thr Leu Leu Phe Tyr Ile Gly Gly His Leu Val Arg Asp  
 597 20 25 30  
 599 Asn Asp His Pro Asp His Ser Ser Arg Glu Leu Ser Lys Ile Leu Ala  
 600 35 40 45  
 602 Lys Leu Glu Arg Leu Lys Gln Gln Asn Glu Asp Leu Arg Arg Met Ala  
 604 50 55 60  
 606 Glu Ser Leu Arg Ile Pro Glu Gly Pro Ile Asp Gln Gly Thr Ala Thr  
 607 65 70 75 80  
 609 Gly Arg Val Arg Val Leu Glu Glu Gln Leu Val Lys Ala Lys Glu Gln  
 610 85 90 95  
 612 Ile Glu Asn Tyr Lys Lys Gln Ala Arg Asn Gly Leu Gly Lys Asp His  
 613 100 105 110  
 615 Glu Ile Leu Arg Arg Arg Ile Glu Asn Gly Ala Lys Glu Leu Trp Phe  
 616 115 120 125  
 618 Phe Leu Gln Ser Glu Leu Lys Lys Leu Lys His Leu Glu Gly Asn Glu  
 619 130 135 140  
 621 Leu Gln Arg His Ala Asp Glu Ile Leu Leu Asp Leu Gly His His Glu  
 622 145 150 155 160  
 624 Arg Ser Ile Met Thr Asp Leu Tyr Tyr Leu Ser Gln Thr Asp Gly Ala  
 625 165 170 175  
 627 Gly Asp Trp Arg Glu Lys Glu Ala Lys Asp Leu Thr Glu Leu Val Gln  
 628 180 185 190  
 630 Arg Arg Ile Thr Tyr Leu Gln Asn Pro Lys Asp Cys Ser Lys Ala Arg  
 631 195 200 205  
 633 Lys Leu Val Cys Asn Ile Asn Lys Gly Cys Gly Tyr Gly Cys Gln Leu  
 634 210 215 220  
 636 His His Val Val Tyr Cys Phe Met Ile Ala Tyr Gly Thr Gln Arg Thr  
 637 225 230 235 240  
 639 Leu Ile Leu Glu Ser Gln Asn Trp Arg Tyr Ala Thr Gly Gly Trp Glu  
 640 245 250 255  
 642 Thr Val Phe Arg Pro Val Ser Glu Thr Cys Thr Asp Arg Ser Gly Leu  
 643 260 265 270  
 645 Ser Thr Gly His Trp Ser Gly Glu Val Asn Asp Lys Asn Ile Gln Val  
 646 275 280 285  
 648 Val Glu Leu Pro Ile Val Asp Ser Leu His Pro Arg Pro Pro Tyr Leu  
 649 290 295 300  
 651 Pro Leu Ala Val Pro Glu Asp Leu Ala Asp Arg Leu Leu Arg Val His  
 652 305 310 315 320  
 654 Gly Asp Pro Ala Val Trp Trp Val Ser Gln Phe Val Lys Tyr Leu Ile  
 655 325 330 335

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/575,261

DATE: 04/24/2006

TIME: 16:15:41

Input Set : A:\seq list.txt

Output Set: N:\CRF4\04242006\J575261.raw

```

657 Arg Pro Gln Pro Trp Leu Glu Lys Glu Ile Glu Glu Ala Thr Lys Lys
658          340          345          350
660 Leu Gly Phe Lys His Pro Val Ile Gly Val His Val Arg Arg Thr Asp
661          355          360          365
663 Lys Val Gly Thr Glu Ala Ala Phe His Pro Ile Glu Glu Tyr Met Val
664          370          375          380
666 His Val Glu Glu His Phe Gln Leu Leu Ala Arg Arg Met Gln Val Asp
667 385          390          395          400
669 Lys Lys Arg Val Tyr Leu Ala Thr Asp Asp Pro Thr Leu Leu Lys Glu
670          405          410          415
672 Ala Lys Thr Lys Tyr Ser Asn Tyr Glu Phe Ile Ser Asp Asn Ser Ile
673          420          425          430
675 Ser Trp Ser Ala Gly Leu His Asn Arg Tyr Thr Glu Asn Ser Leu Arg
676          435          440          445
678 Gly Val Ile Leu Asp Ile His Phe Leu Ser Gln Ala Asp Phe Leu Val
679          450          455          460
681 Cys Thr Phe Ser Ser Gln Val Cys Arg Val Ala Tyr Glu Ile Met Gln
682 465          470          475          480
684 Thr Leu His Pro Asp Ala Ser Ala Asn Phe His Ser Leu Asp Asp Ile
685          485          490          495
687 Tyr Tyr Phe Gly Gly Gln Asn Ala His Asn Gln Ile Ala Val Tyr Pro
688          500          505          510
690 His Lys Pro Arg Thr Glu Glu Glu Ile Pro Met Glu Pro Gly Asp Ile
691          515          520          525
693 Ile Gly Val Ala Gly Asn His Trp Asp Gly Tyr Ser Lys Gly Ile Asn
694          530          535          540
696 Arg Lys Leu Gly Lys Thr Gly Leu Tyr Pro Ser Tyr Lys Val Arg Glu
697 545          550          555          560
699 Lys Ile Glu Thr Val Lys Tyr Pro Thr Tyr Pro Glu Ala Glu Lys
E--> 700          565          570
2297 <210> SEQ ID NO: 64
2298 <211> LENGTH: 235
2299 <212> TYPE: PRT
2300 <213> ORGANISM: Homo sapiens
2302 <400> SEQUENCE: 64
2303 Leu Pro Ala Gln Val Ala Phe Thr Pro Tyr Ala Pro Glu Pro Gly Ser
2304 1          5          10          15
2306 Thr Cys Arg Leu Arg Glu Tyr Tyr Asp Gln Thr Ala Gln Met Cys Cys
2307          20          25          30
2309 Ser Lys Cys Ser Pro Gly Gln His Ala Lys Val Phe Cys Thr Lys Thr
2310          35          40          45
2312 Ser Asp Thr Val Cys Asp Ser Cys Glu Asp Ser Thr Tyr Thr Gln Leu
2313          50          55          60
2315 Trp Asn Trp Val Pro Glu Cys Leu Ser Cys Gly Ser Arg Cys Ser Ser
2316 65          70          75          80
2318 Asp Gln Val Glu Thr Gln Ala Cys Thr Arg Glu Gln Asn Arg Ile Cys
2319          85          90          95
2321 Thr Cys Arg Pro Gly Trp Tyr Cys Ala Leu Ser Lys Gln Glu Gly Cys
2322          100          105          110

```

## RAW SEQUENCE LISTING

DATE: 04/24/2006

PATENT APPLICATION: US/10/575,261

TIME: 16:15:41

Input Set : A:\seq list.txt

Output Set: N:\CRF4\04242006\J575261.raw

```

2324 Arg Leu Cys Ala Pro Leu Arg Lys Cys Arg Pro Gly Phe Gly Val Ala
2325      115      120      125
2327 Arg Pro Gly Thr Glu Thr Ser Asp Val Val Cys Lys Pro Cys Ala Pro
2328      130      135      140
2330 Gly Thr Phe Ser Asn Thr Thr Ser Ser Thr Asp Ile Cys Arg Pro His
2331 145      150      155      160
2333 Gln Ile Cys Asn Val Val Ala Ile Pro Gly Asn Ala Ser Met Asp Ala
2334      165      170      175
2336 Val Cys Thr Ser Thr Ser Pro Thr Arg Ser Met Ala Pro Gly Ala Val
2337      180      185      190
2339 His Leu Pro Gln Pro Val Ser Thr Arg Ser Gln His Thr Gln Pro Thr
2340      195      200      205
2342 Pro Glu Pro Ser Thr Ala Pro Ser Thr Ser Phe Leu Leu Pro Met Gly
2343      210      215      220
2345 Pro Ser Pro Pro Ala Glu Gly Ser Thr Gly Asp
E--> 2346 225      230      235

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RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/575,261

DATE: 04/24/2006  
TIME: 16:15:43

Input Set : A:\seq list.txt  
Output Set: N:\CRF4\04242006\J575261.raw

**Invalid Line Length:**

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:17; Line(s) 832  
Seq#:74; Line(s) 2501  
Seq#:75; Line(s) 2558  
Seq#:76; Line(s) 2666



**VERIFICATION SUMMARY**

PATENT APPLICATION: US/10/575,261

DATE: 04/24/2006

TIME: 16:15:43

Input Set : A:\seq list.txt

Output Set: N:\CRF4\04242006\J575261.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No  
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:48 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:52 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:56 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:60 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:64 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:68 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:72 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:76 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:80 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:84 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:88 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:92 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:97 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:101 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:105 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:109 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:113 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:117 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:121 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1  
L:153 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:2  
M:332 Repeated in SeqNo=2  
L:208 M:252 E: No. of Seq. differs, <211> LENGTH:Input:376 Found:372 SEQ:2  
L:584 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7  
L:700 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:8  
L:1396 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:46  
L:1529 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:48  
L:1676 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:51  
L:1786 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:53  
L:2346 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:64